

1 ATAAGAATGC GGCCGCATGA CCCCGCAGCT TCTCCTGGCC CTTGTCCTCT
 51 GGGCCAGCTG CCCGCCCCGC AGTGGAAGGA AAGGGCCCCC AGCAGCTCTG
 101 AACTGCCCC GGGTGCAATG CCGAGCCTCT CGGTACCCGA TCGCCGTGGA
 151 TTGCTCCTGG ACCCTGCCGC ATGACCCCGC AGCTTCTCCT GGCCCTTGTC
 201 CTCTGGGCCA GCTGCCCCGCC CTGCGGTGGA AGGAAAGGGC CCCAGCAGC
 251 TCTGACACTG CCCCGGGTGC AATGCCGAGC CTCTCGGTAC CCGATCGCCG
 301 TGGATTGCTC CTGGACCCTG CCGCCTGCTC CAAACTCCAC CAGCCCCGTG
 351 TCCTTCATTG CCACGTACAG GCTCGGCATG GCTGCCCCGG GCCACAGCTG
 401 GCCTTGCCCTG CAGCAGACGC CAACGTCCAC CAGCTGCACC ATCACGGATG
 451 TCCAGCTGTT CTCCATGGCT CCCTACGTGC TCAATGTCAC CGCCGTCCGC
 501 CCCTGGGGCT CCAGCAGCAG CTTCGTGCCT TTCATAACAG AGCACATCAT
 551 CAAGCCCGAC CCTCCAGAAG GCGTGCGCCT AAGCCCCCTC GCTGAGCGCC
 601 AGCTACAGGT GCAGTGGGAG CCTCCCGGGT CCTGGCCCTT CCCAGAGATC
 651 TTCTCACTGA AGTACTGGAT CCGTTACAAG CGTCAGGGAG CTGCGCGCTT
 701 CCACCGGGTG GGGCCCATTG AAGCCACGTC CTTCATCCTC AGGGCTGTGC
 751 GGCCCCGAGC CAGGTACTAC GTCCAAGTGG CGGCTCAGGA CCTCACAGAC
 801 TACGGGGAAC TGAGTGAAGT GAGTCTCCCC GCCACTGCCA CAATGAGCCT
 851 GGGCAAGTAG ACTAGTCC

Figure 1

1 MTPQLLLALV LWASCPPRSG RKGPPAALTl PRVQCRASRY PIAVDCSWTL
51 PHDPAASPGP CPLGQLPALR WKERAPSSSD TAPGAMPSLS VPDRRGLLLD
101 PAACSKLHQP RVLHCHVQAR HGCPGPQLAL PAADANVHQL HHHGCPAVLH
151 GSLRAQCHRR PPLGLQQQLR AFHNRAHHQA RPSRRRAPKP PR*

Figure 2

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EBI-3 (AAA93193)	MTPQLLLALVLWASCPPCSGRKGPPAALTLP RVQCRASRYPIAVDCSWTLPPAPNSTSPV	60
EBI-3-alt		
	MTPQLLLALVLWASCPPRSGRKGPPAALTLP RVQCRASRYPIAVDCSWTLPHDPAASGPGP	
EBI-3 (AAA93193)	SFIATYRLGMAARGHSWPCLQQTPTSTSTCTITDVQLFSMAPYVLNVTAVHPWGSSSSSFPV	120
EBI-3-alt	CPLGQLPALRWKERAPSSSDTAPGAMPSLSVPDRRGLLLDPAACSKLHQPRVLHCHVQAR	
EBI-3 (AAA93193)	FITEHI IKPDPPEGVRLSPLAERHVQVQWEPPGSWPFPEIFSLKYWIRYKRQGAARFHRV	180
EBI-3-alt	HGCPGPQLALPAADANVHQLHHHGCPAVLHGSLRAQCHRRPPLGLQQQLRAFHNRAHHQA	
EBI-3 (AAA93193)	GPIEATSFILRAVRPRARYYVQVAAQDLTDYGELSDWSLPATATMSLGK	
EBI-3-alt	RPSRRRAPKPPR*	

Figure 4